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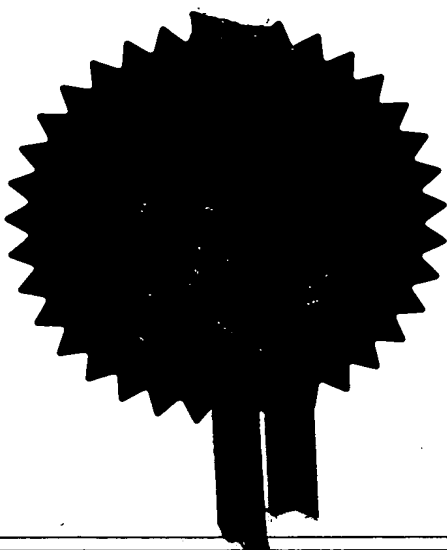
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Recombinant microorganisms

- The present invention relates to recombinant microorganisms, in particular gut-colonising organisms, which are useful for example in the delivery of antigenic material and thus form the basis of vaccines. Vaccines comprising these organisms and promoter sequences for use in them form a further aspect of the invention.
- 10 Attenuated mutants of *Salmonella typhi* (e.g. *aroA*, *aroC*, *htrA*) are currently being evaluated as live, oral vaccines against typhoid fever (Tacket CO, et al., Infect. Immun. 1997;65:452-6). These mutants have also attracted attention as carriers for guest (vaccine) antigens but suitable animal models for testing these vaccines are not available. In view of this, many workers have used *Salmonella typhimurium aroA* expressing guest antigens for investigating the immune responses induced after oral vaccination of mice.
- 20 The unregulated expression of foreign genes within *Salmonella* species such as *S. typhimurium* can lead to plasmid instability, yet the stable expression of the guest antigen at the appropriate site in the body is necessary for the induction of a protective response. One approach to promote the stable expression of guest antigens involves the chromosomal integration of the heterologous gene. However, this may reduce the immune response because of gene dosage effects (Covone M G, et al., Infect. Immun. 1998;66:224-31).
- 30 The balanced lethal system (Curtiss R III, et al., Res. Microbiol. 1990;141:797-805, Nakayama K, et al., Bio/Technology 1988;6:693-97) relies on the complementation of a lethal mutation by a plasmid which also encodes the guest antigen. Whilst this ensures retention of the plasmid, the gene encoding the guest antigen itself may be deleted. An alternative approach involves the use of promoters which are induced within
- 35
-
-

host tissues to direct guest antigen expression at that site. Because the gene is only expressed after certain environmental cues have been recognised, this approach might reduce the selective pressure towards deleting the gene.

5

This solution to the problem of expression of guest antigens has also been identified by other workers. A variety of antigens have been expressed in *S. typhimurium* from the *nirB* promoter which is upregulated under anaerobic conditions and within host
10 cells (Oxer M D, et al. Nucleic Acids Res. 1991;19:2889-92). Guest antigens delivered using the *nirB* promoter system induce superior responses than the same antigens delivered from a constitutive promoter. In addition, the *nirB* promoter-driven genes were maintained more effectively in the *Salmonella* host
15 strain. More recently, it has been shown that the *htrA* and *osmC* promoter can be used to direct expression of guest antigens in *Salmonella* (McSorley S J, et al., Infect. Immun. 1997;65:171-78, Roberts M, et al., Infect. Immun. 1998;66:3080-87). However, it is likely that these promoters will not be suited to the
20 expression of all guest antigens.

Immunisation with the F1-antigen of *Y. pestis* has previously been shown to induce an antibody-mediated protective response against plague (Green M, et al., FEMS Microbiology and
25 Immunology, 1998;23:107-13) and we have previously shown that the F1-antigen can be expressed in *S. typhimurium* (Oyston P C F, et al., Infect. Immun. 1995;63:563-68, Titball R W, et al., Infect. Immun. 1997;65:1926-30). The antigenic properties of F1-antigen have been exploited to investigate the ways in which
30 different promoters, which are induced at different sites in the body, can be used to induce different antibody responses to guest antigens expressed in *S. typhimurium*. It is known that the invasion and spread of *S. typhimurium* within the host is accompanied by the expression of different subsets of genes
35 which are involved in processes such as attachment and invasion,

penetration of the epithelium and the infection of deep lymphoid tissue.

The OmpR/EnvZ two component regulatory system responds to changes in the osmotic strength and pH within *S. typhimurium* (Foster J W, et al., Microbiology 1994;140:341-52). It has been suggested that this system might play a role in allowing the bacterium to survive in the gut by regulating the expression of outer membrane porins such as OmpC (Pratt L A, et al., American Society for Microbiology, ASM Press, Washington DC, 1995, pp105-27, Nikaido H, et al., Cellular and Molecular Biology. American Society for Microbiology, Washington DC. 1987, pp7-22, García Vescovi E. et al., Cell. 1996;84:165-74).

The PhoP/PhoQ two-component regulatory system controls virulence properties such as survival within macrophages, resistance to host defence antimicrobial peptides and acid pH, invasion of epithelial cells, the formation of spacious vacuoles and the processing and presentation of antigens by activated macrophages (Miller S I. et al., Proc. Natl. Acad, Sci USA 1989;86:5054-58, Fields P I, et al., Science 1989;243:1059-62, Pegues D A, et al., Mol. Microbiol. 1995;17:169-81, Wick M J, et al., Mol. Microbiol. 1995;16:465-76), in response to environmental magnesium concentration (García Vescovi E. et al., Cell. 1996;84:165-74). Over forty genes are regulated by this system in *S. typhimurium* (Soncini F C, et al., J. Bacteriol. 1996;178:5092-99) including the *phoP* gene, which is autoregulated (Soncini F C, et al., J. Bacteriol. 1995;177:4364-71) and the *pagC* gene which encodes an envelope protein required for survival in the macrophage (Alpuche-Aranda C M, et al., Proc. Natl. Acad. Sci. USA 1992;89:10079-83).

The present invention provides a recombinant gut-colonising microorganism which comprises a promoter selected from the P_{ompC} , P_{phoP} and P_{pagC} or fragments or variants thereof which can act as promoters, said promoter being operatively interconnected with a

nucleic acid which encodes a heterologous protein, able to induce a protective immune response against a different organism, in a mammal to which it is administered. The term "heterologous protein" refers to proteins which are not native to the microorganism strain.

The three promoters (P_{phoP} , P_{pagC} and P_{ompC} ,) which are included in the organisms of the invention are induced at different stages in the infection process, and hence at different sites in the body. This approach allows the induction of different immune responses which provide protection against pathogens which colonise different host cell compartments. The sequence of these promoters has been elucidated previously, and these are given hereinafter in Figure 6 as SEQ ID NOS 2, 3 and 4 respectively.

Their expression has been compared to that of the constitutively expressed *lacZ* gene promoter. As a result, recombinant gut-colonising microorganisms wherein antigen expression is driven by P_{phoP} promoter forms a preferred embodiment of the invention.

The development of effective vaccines against pathogens is dependent not only on the identification of the appropriate protective antigens but also on the induction of an immune response at the site in the body which provides maximum protection against disease. For some pathogens, serum antibody provides protection against disease. However, many pathogens enter the body at a mucosal surface and protection against these diseases might therefore be dependent on the induction of mucosal immune responses. The *Salmonella* vaccine vector system is ideally suited to the delivery of many vaccine antigens since the vaccine delivery mechanism accurately mimics the natural disease, entering the body via the gut.

Thus in particular embodiment, the recombinant gut-colonising microorganism comprises a *Salmonella* spp. such as *Salmonella typhimurium* or *Salmonella typhi*.

- 5 Recombinant gut-colonising microorganisms of the invention are suitably attenuated so that the host does not experience significant harmful effects as a result of infection by the microorganism. Examples of attenuated mutants include *aro* mutants such as *aroA* and *aroC* mutants, apartate β -semi-aldehyde
 10 dehydrogenase (ASD) mutants, purine biosynthesis mutants, branched chain amino acid biosynthesis mutants, galactose epimerase (*galE*) mutants, regulatory mutants such as *phoP* and *phoQ* mutants, *htrA* serine protease mutants and adenyl cyclase mutants. Particular attenuated strains of *Salmonella*, such as
 15 *Salmonella typhi* include *aroA*, *aroC* and *htrA* mutants or triple mutants including all three mutations.

- Recombinant gut-colonising microorganism as described above can be used to deliver a variety of antigenic agents which can be
 20 used to induce a protective immune response against a wide range of pathogens. Pathogens which may be targeted in this way are those of humans or animals and include those listed in the Health and Safety Executive: "Categorisation of Biological Agents according to Hazard and Category of Containment", HMSO,
 25 ISBN 0717610381. Particular examples of antigenic agents which may be included in the recombinant organisms of the invention include those protective against tetanus such as tetanus toxin H_c fragment, those protective against Botulinum such as *botulinum* toxin H_c fragment, those protective against *Bacillus anthracis*
 30 such as *Bacillus anthracis* protective antigen (PA), those protective against *Bordatella pertussis* such as *Bordatella pertussis* P69 antigen, those protective against *Schistoma mansonii* such as *Schistoma mansonii* glutathione-S-transferase, those protective against cholera such as Fibrio cholera β sub-
 35 unit, those protective against Herpes simplex virus (HSV) such as HSV glycoprotein D, those protective against HIV infection such

as HIV envelope protein, and those protective against *Escherischia coli* such as *E. coli* LTB subunit or *E. coli* K88 antigen. In particular, it has been found that where the heterologous protein, is able to induce a protective immune response against *Yersinia pestis*, useful protective immunity is found. Examples of antigens which can produce such as response include the F1-antigen of *Yersinia pestis* or an antigenic fragment or variant thereof, or the V-antigen of *Yersinia pestis* or combinations thereof as described in WO 96/28551.

10

The expression "variant" refers to sequences of amino acids which differ from the base sequence from which they are derived in that one or more amino acids within the sequence are substituted for other amino acids. Amino acid substitutions may be regarded as "conservative" where an amino acid is replaced with a different amino acid with broadly similar properties. Non-conservative substitutions are where amino acids are replaced with amino acids of a different type. Broadly speaking, fewer non-conservative substitutions will be possible without altering the biological activity of the polypeptide. Suitably variants will be at least 60% homologous, preferably at least 75% homologous, and more preferably at least 90% homologous to the base sequence. Homology in this instance can be determined using in particular the Needleman-Wunsch algorithm with gap penalty of 8 using a standard PAM scoring matrix (Needleman S.B. and Wunsch C.D., J. Mol Biol. 1970, vol 48, 443-453).

The recombinant gut-colonising microorganisms described above are thus particularly suitable for use in the preparation of vaccines for therapeutic or prophylactic purposes, where they may be combined with a pharmaceutically acceptable carrier or diluent, as would be understood in the art.

In particular, the vaccines will be formulated so that they are adapted for oral administration and that the microorganism

remains viable throughout any storage period. Thus they may preferably be in a form liquid form such as aqueous or oily suspensions, emulsions, syrups or elixirs.

- 5 The size of the dose for therapeutic or prophylactic purposes of will vary according to a wide variety of factors including the nature of the protective immune response sought, the nature of the antigen being employed, the severity of the conditions, the dosage regime in terms of primary and secondary boosting, the
10 age and sex of the animal or patient and the gut-colonising ability of the particular microorganism used. In general however, a dosage of microorganism in the range of from 10^6 to 10^9 cfu will be administered as a single dosage.
- 15 Vaccine compositions may further comprise a buffer such as a bicarbonate buffer, in order to neutralise stomach acid.

Thus in a further aspect, the invention provides a method of inducing a protective immune response against a pathogen in a
20 mammal, said method comprising administering to said mammal a recombinant gut-colonising microorganism which comprises a promoter selected from the P_{ompC} , P_{phoP} and P_{pagC} or fragments or variants thereof which can act as promoters, said promoter being operatively interconnected with a nucleic acid which encodes an
25 antigen protein, able to induce a protective immune response against said pathogen, in a mammal to which it is administered.

In yet a further aspect, the invention provides the use of a promoter selected from P_{ompC} , P_{phoP} and P_{pagC} in the production of a
30 vaccine comprising a recombinant gut-colonising organism.

The promoters used in this study are induced at specific sites in the body. They are preferably cloned into the microorganism in a low copy number vector, because high copy number plasmids
35 have been shown to be unstable in *S typhimurium* (Coulson N M, et al., Microb Pathog. 1994;16:305-11).

The PhoP gene would be expected to be expressed at a basal level from the *PhoPp2* promoter and upregulated in the phagosome of host cells as a result of activation of the *PhoPp1* promoter (Soncini F C, et al., J. Bacteriol. 1995;177:4364-71). The PhoP / PhoQ regulatory system has been shown to regulate the expression of a variety of genes including *pagC*, and to be important for survival in macrophages (Miller S I. et al., Proc. Natl. Acad. Sci USA 1989;86:5054-58, Wick M J, et al., Mol. Microbiol. 1995;16:465-76).

Genes regulated by the PhoP/PhoQ system are also important for the virulence of orally delivered bacteria (Galan J E, et al., Microb Pathog. 1989;6:433-43). Expression of the *ompC* is gene is upregulated under conditions of high osmotic strength (Foster J W, et al., Microbiology 1994;140:341-52, Nikaido H, et al., Cellular and Molecular Biology. American Society for Microbiology, Washington DC. 1987, pp7-22), such as those found within the gut, under control of the OmpR/EnvZ regulatory system (Pratt L A, et al., American Society for Microbiology, ASM Press, Washington DC, 1995, pp105-27).

Whilst the different plasmids in *S. typhimurium* SL3261 were stable in vitro, there were marked differences in the stability of the plasmids in bacteria which had been delivered to mice by the oral route. Bacteria expressing the F1-antigen from the *PagC* promoter showed a much reduced ability to colonise mesenteric lymph nodes and appeared incapable of further invasion of the host. It is possible that the additional copies of the *pagC* promoter and upstream regulatory regions titrated out the available PhoP activator within the cell, and that this prevented the bacterium from responding to the environmental changes encountered after uptake by M-cells. However, recombinant *Salmonella* containing the *P_{phoP}*-F1 plasmid did not show a similar inability to invade the host.

This finding might be in accordance with the suggestion that *phoP* expression is only partially autoregulated by the *phoP* gene product (Fields P I, et al., Science 1989;243:1059-62).

5 Additionally, it is possible that the high level of expression of F1-antigen from the *pagC* promoter in vivo placed a lethal metabolic load on the host bacterium.

These promoters are regulated by a variety of environmental stimuli in a manner which is not fully defined. Therefore, it
10 is difficult to make meaningful comparisons of the strengths of these promoters in vitro. Thus, in vivo testing of these promoters to identify those most suitable for use for the expression of guest antigens has been carried out.

15 All of the recombinant *Salmonella* induced similar levels of antibody against the whole bacterium. This finding was unexpected for bacteria containing *pP_{pagC}*-F1, since these bacteria were unable to invade deep host tissues and were recovered only at low levels from mesenteric lymph nodes. This recombinant
20 *Salmonella* also induced IgG and IgA antibody against the F1-antigen. This suggests that the initial interaction of the bacteria with M-cells is critical in determining the immune response to the bacterium and to guest antigens. This conclusion is supported by the finding that *Salmonella*
25 containing *pP_{pagC}*-F1 induced mucosal antibody to the F1-antigen whereas bacteria expressing the F1-antigen expressed from the *lacZ* or *ompC* promoters failed to induce mucosal responses. Therefore, the measurement of the colonisation of spleen or liver tissues, as an indicator of vaccine potential of
30 recombinant *Salmonella*, may not always be useful.

Similar conclusions were reached by Covone et al. (Covone M G, et al., Infect. Immun. 1998;66:224-31) who showed that effective delivery of the LTK63 guest antigen to the immune system was
35 effective only when the antigen was delivered during the early stages of invasion and by McSorley et al. (McSorley S J, et al.,

Infect. Immun. 1997;65:171-78) who showed that recombinant *Salmonella* expressing glycoprotein 63 from the *osmC* promoter were unable to invade tissue beyond the mediastinal lymph nodes, yet induced protection against *Leishmania major*. This might
5 also explain why killed *Salmonella* with or without guest antigens, which are clearly not able to invade deep host tissues, are able to induce an immune response (Thatte J, et al., Int. Immunol. 1993;5:1431-36).

- 10 The ability of *Salmonella* expressing P_{phoP} -F1 to induce mucosal antibody responses to the F1-antigen in both the gut and the lungs, whereas a constitutive promoter (P_{lacZ}) failed to induce such responses clearly demonstrates the utility of *in vivo* induced promoters for the induction of appropriate antibody
15 responses. This promoter system will be particularly useful for other applications where a mucosal antibody response is important for protection against disease.

The invention will now be particularly described by way of
20 Example with reference to the accompanying diagrammatic drawings in which:

Figure 1 is a plasmid diagram illustrating plasmids used in the preparation of microorganisms in accordance with the invention;
25

Figure 2 is a graph showing the levels of colonisation of spleen tissues of mice, 11 days after dosing with recombinant microorganisms of the invention;

30 Figure 3 shows graphs illustrating IgG serum antibody levels in mice to the carrier bacterium, (Fig 3a) and to the F1 antigen (Fig 3b), 21, 28 and 98 days after immunisation;

Figure 4 is a graph showing the isotype of the F1 antibody found
35 in mice serum on day 98, where the blank column represents the

amount of the IgG1_a? and the shaded column represents the IgG2_a isotype in all groups of immunised animals:

Figure 5 is a graph showing the levels of circulating IgA antibody to F1-antigen or the levels of IgA antibody to F1-antigen in gut (blank column) or lung (shaded column) wash samples; and

Figure 6 shows sequences of promoters used in the evaluation of the invention.

Example 1

Preparation of Bacterial strains, cultivation and enzymes

Escherichia coli strain JM109 and *S. typhimurium* strains LB5010 (rm⁺ *galE*), SL3261 (*aroA*) or SL1344 (a mouse-virulent strain; (Zhang X, et al., Infect. Immun. 1997;65:5381-7) were cultured on L-agar or in L-broth, supplemented with ampicillin (05µg/ml) where appropriate. Enzymes used for DNA cloning and amplification procedures were obtained from BCL limited (Lewes, Sussex, UK). PCR reactions were carried out using a Perkin Elmer 9600 (P.E. Applied Biosystems, Warrington, UK) thermal cycler with cycle conditions of 95°C, 5 min, followed by 50 cycles of 95°C, 5s; 45°C, 5s; 72°C, 5s, followed by 10 min at 72°C.

Plasmids containing promoters for expression of F1-antigen were then produced. The promoters for the *phoP*, *pagC* and *ompC* genes have previously been mapped and upstream regulatory regions identified (Soncini F C, et al., J. Bacteriol. 1995;177:4364-71, Pulkkinen W S, et al., J. Bacteriol. 1991;173:86-9, Puente J L, et al., Gene. 1987;61:75-83, Puente J L, et al., Gene. 1989;83:197-206). For the *phoP* gene promoter a 139bp DNA fragment was identified which included the *phoPp1* and *phoPp2* gene promoters and 80 bp upstream of the -35 site which has been predicted to form step loop structures (Soncini F C, et al., J. Bacteriol. 1995;177:4364-71). For the *pagC* gene promote a 715

bp DNA fragment included 125 bp upstream of the -35 region (Pulkkinen W S, et al., J. Bacteriol. 1991;173:86-9). For the *ompC* gene promoter a 371 bp DNA fragment included a 275 bp region upstream of the -35 region (Puente J L, et al., Gene. 1987;61:75-83, Puente J L, et al., Gene. 1989;83:197-206).

These DNA fragments were amplified from *S. typhimurium* strain SL1344 genomic DNA using the PCR. For comparison with a constitutive gene promoter, a 196 bp DNA fragment encoding the *lacZ* gene promoter and 140 bp upstream of the -35 region was identified. The 3' end of all of the DNA fragments terminated before the SD regions associated with the genes. These promoters were cloned upstream of the *cafI* open reading frame (encoding the *Y. pestis* F1-antigen) and SD region in a low copy number vector (pBR322; Fig 1) and the recombinant plasmids (pP_{ompC}-F1, pP_{p_hoP}-F1, pP_{p_agC}-F1 or pP_{lacZ}-F1) transformed into *S. typhimurium* SL3261 (*aroA*).

Oligonucleotide primers were designed to amplify promoter regions using the PCR (Table 1).

Table 1

Oligonucleotide pair	SEQ ID No	Prom
5' AAGGAAAAAAGCGGCCGCGCCCAATACGCAAACCG 3'	5	
5' GAATTC <u>ACTAGT</u> ATTGTTATCCGCGCTCACAAT 3'	6	<i>Plac</i>
5' AAGGAAAAAAGCGGCCGCTGACTCTGGTCGACGAACTTA 3'	7	
5' CTAGTCTAGATGTGTTAACCAATAAGAACAGTCTA 3'	8	<i>PphoP</i>
5' AAGGAAAAAAGCGGCCGCTAAACAGACATTCAGAAGTGAATG3'	9	
5' CTAGTCTAGAAATATGCCTTTATTGCTTTTTTATG 3'	10	<i>PompC</i>
5' AAGGAAAAAAGCGGCCGCGTTAACCACTCTTAATAATAATG 3'	11	
5' CTAGACTAGTTACTACTTATTATTACG 3'	12	<i>PpagC</i>

Restriction sites are shown underlined.

The primers included unique *Not*I, *Xba*I or *Spe*I sites. Regions amplified included the -10 and -35 regions and upstream regulatory binding sites, but excluded the Shine-Dalgarno (SD) ribosome binding site.

5

After PCR amplification of the promoter regions from *S.*

typhimurium SL 1344 template DNA (or plasmid pUC18 template DNA for amplification of the *lac* promoter), the DNA fragments were purified using Microcon 100 centrifugal concentrations

10 (Millipore, Watford, UK). The purified DNA fragments were

cloned into suitable digested plasmid pBluescript SK-,

electroporated into *E. coli* JM101 and the cloned fragments were nucleotide sequenced to ensure their authenticity. After

15 digestion of the recombinant plasmids with *Sac*I and *Bss*HI and agarose gel electrophoresis, DNA fragments containing the

promoter regions were purified using Qiaex (Qiagen Ltd, Crawley, UK) and blunt ended using Klenow fragment (Sambrook J, Fritsch E F, Maniatis T. 1989. Molecular Cloning: A Laboratory Manual, 2nd ed. Cold Spring Harbor Laboratory press, New York).

20

The authentic promoter sequences were then cloned into plasmid pBR322 which had been digested with *Eco*R1 and *Nru*I and then blunt ended using Klenow fragment (Sambrook J, Fritsch E F, Maniatis T. 1989. Molecular Cloning: A Laboratory Manual, 2nd ed.

25 Cold Spring Harbor Laboratory press, New York). The orientation of the cloned DNA fragment in the plasmid was determined by analysing, using agarose gel electrophoresis, the DNA fragments obtained after digestion with *Xba*I, *Ssp*I or *Sty*I.

30 A DNA fragment which encoded the *Caf*I open reading frame and the ribosome binding site was isolated after digestion of plasmid pORF1 (Oyston P C F, et al., Infect. Immun. 1995;63:563-68) with *Eco*R1 followed by blunt ending of the DNA and further digestion with *Hind*III. The purified DNA fragment was ligated with

35 promoter plasmids with which had been digested with *Sma*I and

HindIII. The final recombinant plasmids were transformed into *E. coli* strain JM109.

Plasmids were isolated from *E. coli* (Sambrook J, Fritsch E F, Maniatis T. 1989. Molecular Cloning: A Laboratory Manual, 2nd ed. Cold Spring Harbor Laboratory press, New York) and electroporated into *S. typhimurium* SL3281 (*aroA*) after passage through *S. typhimurium* LB5010 to ensure methylation of the DNA.

10 Example 2

The stability of the different plasmids encoding F1-antigen driven from different promoters in *S. typhimurium* SL3261 was determined after culture of the bacteria in L-broth for 24hr (*in vitro* stability) and enumeration of bacteria which grew on L-agar or L-agar containing ampicillin (Sambrook J, Fritsch E F, Maniatis T. 1989. Molecular Cloning: A Laboratory Manual, 2nd ed. Cold Spring Harbor Laboratory press, New York). The results indicated that all of the plasmids were retained by at least 80% of the bacteria which had been cultured *in vitro* (pP_{ompC}-F1, 83%; pP_{phoP}-F1, 100%; pP_{pagC}-F1, 98%; pP_{lacZ}-F1, 95%).

Example 3

Colonisation of host tissues

The *in vivo* stability of plasmids was determined by inoculating groups of 10 female BALB/c mice orally with 10⁹ cfu of bacteria in 100µl of PBS, and enumerating bacteria isolated 11 days later from homogenised spleen tissue on L-agar or L-agar + ampicillin.

In the case of the P_{pagC}-F1 construct and the *S. typhimurium* SL3261 control strain, bacteria were also cultured from mesenteric lymph nodes (10/mouse) each homogenised in 2ml of PBS or from homogenised liver tissue which were removed on 11 days after dosing. Bacteria were enumerated as described above.

With the exception of the bacteria containing the P_{pagC} -F1 plasmid and the SL3261 control bacteria, ampicillin-resistant bacteria could be recovered from all of the spleens isolated from orally dosed mice. The *in vivo* stability of all of the plasmids within
5 *Salmonella* was lower than the stability of the plasmids within *Salmonella* cultured *in vitro*.

Although mice were dosed orally with similar numbers of bacteria there were marked differences in the level of colonisation of
10 spleen tissues at day 11 (Fig 2). The highest levels of colonisation were found with bacteria containing pP_{phoP} -F1. Low numbers of *Salmonella* containing pP_{pagC} -F1 were recovered from spleen tissues (maximum 50 cfu/spleen) and none of the bacterial colonies were ampicillin resistant. To investigate why the
15 presence of the pP_{pagC} -F1 affected the colonising ability of this *Salmonella*, a more detailed study was undertaken which involved the enumeration of bacteria in the liver and in mesenteric lymph nodes 11 days after oral dosing. The levels of SL3261 containing pP_{pagC} -F1 isolated from liver tissues were similar to
20 those isolated from spleens (data not shown). Whereas the mean number of SL3261 isolated from mesenteric lymph nodes was 2.2×10^3 cfu, SL3261 containing pP_{pagC} -F1 could be isolated only at low level (mean 20 cfu) from these tissues.

25 Example 4

Serum antibody responses to F1-antigen

Groups of 3 or 8 female BALB/c mice were immunised via intragastric intubation on days 0 and 14 with 1×10^9 cells of the P_{ompC} -F1, P_{phoP} -F1, P_{pagC} -F1 or P_{lacZ} -F1 constructs, or the
30 control *S. typhimurium* strain, SL3261, in 0.1 ml of phosphate-buffered saline (PBS). Bacteria were grown statically overnight at 37°C. All oral inoculations were carried out with a stainless steel gavage needle without an anaesthetic. The inoculum dose was verified by plating serial dilutions of each culture on L-
35 agar plates with or without ampicillin.

On days 21, 28 and 98 mice were anaesthetized by intraperitoneal (i.p.) administration of a cocktail of domitor (6 mg per dose) and Ketalar (27mg per dose) and blood was collected by cardiac puncture. Mice were then sacrificed by cervical dislocation.

- 5 Blood was allowed to clot at 4°C overnight prior to centrifugation (10,000 x g, 10 min, 4°C) and the serum stored at -20°C until tested.

- 10 After i.g. dosing with the recombinant *Salmonella*, mice in all groups developed IgG serum antibody to the carrier bacterium, which reached a maximum level 98 days after immunisation (Fig 3a). In the case of SL3261 expressing P_{phoP}-F1, the onset of the immune response was delayed. All groups of mice developed IgG serum antibody to F1-antigen (Fig 3b), which had reached a peak
15 at day 28 and declined by day 98. There was no significant difference in the peak titres to F1-antigen induced by the different recombinant *Salmonella*.

- When the isotype of this antibody was determined on day 98, it
20 was found to be predominantly of an IgG2_a isotype in all groups of immunised animals (Fig 4), suggesting that a Th1-type response was induced. Several other workers have shown that recombinant *S. typhimurium* induce a Th1-type response to the guest antigen (Brett S J, et al., Immunology 1993;80:306-12,
25 Thatte J, et al., Int. Immunol. 1993;5:1431-36).

Example 5

Mucosal antibody responses to F1-antigen

- The ability of the different recombinant *Salmonella* to induce a
30 mucosal antibody response after i.g. dosing was determined by measuring the levels of circulating IgA antibody to F1-antigen or the levels of IgA antibody to F1-antigen in gut or lung wash samples. After dosing as described in Example 4, on days 21 and 28, gut and lung wash samples were collected. Briefly, gut
35 wash samples were collected by resecting a 10 cm length of small intestine and flushing with 5 ml of PBS. Samples were sonicated

for 0.5 min prior to centrifugation (12,000 x g, 30 mins 4°C) and the supernatant was decanted and lyophilized. Broncho-alveolar washings were collected from individual animals by injecting 5 ml of chilled lavage medium (0.9% (w/v) NaCl, 0.05% (v/v) tween 20, 0.1% (w/v) NaN₃ and 1mM phenylmethylsulfonyl fluoride) into the trachea using an intravenous canula and inflating the lungs. A syringe was used to remove the washings, which were subsequently centrifuged (12,000 x g, 30 min, 4°C) prior to lyophilisation of the supernatant fluid. Gut and lung wash samples were reconstituted in 200µl sterile water immediately before use.

All measurements of antibody levels in individual animals were determined in duplicate. For enzyme-linked immunosorbant assays (ELISAs) to determine IgG and IgA titres, 96-well microtiter plates were coated overnight at 4°C either with 50µl 5µg/ml purified F1-antigen (Miller J, et al., FEMS Microbiology and Immunology 1998;21:213-21) in PBS or with 50µl 6µg/ml *S. typhimurium* SL3261 lysate in PBS, prepared as follows. Bacteria were grown statically overnight at 37°C, prior to harvesting and resuspension in PBS to an approximate concentration of 1 x 10¹⁰ cfu/ml. Cells were heat-killed in a boiling water bath for 30 min, cooled on ice and then sonicated on ice for 6 pulses of 30 s. Total protein concentration was determined by a BCA protein assay (Pierce and Warriner, Chester, UK). Plates were blocked for 1 h at 37°C with PBS containing 1% (w/v) skimmed milk powder (BLOTTO). Serum, gut and lung wash samples were diluted in BLOTTO and 50µl volumes were assayed in duplicate in a series of twofold dilutions. After incubation overnight at 4°C, plates were washed three times in PBS with 0.02% (v/v) tween 20. Peroxidase-conjugated secondary antibodies against mouse IgG or IgA (Harlan Sera-Lab Ltd, Loughborough, UK), diluted 1:2000 in BLOTTO were incubated for 1 h at 37°C. The plate was washed as previously and 100µl of 2,2'-azino bis(3-ethylbenzthiazoline-6-

sulfonic acid) substrate (ABTS; Sigma, Poole, UK) was added. Antibody titre was estimated as the maximum dilution of serum giving an absorbance_{414nm} reading 0.1 U above background.

- 5 To determine IgG₁, or IgG_{2a} concentrations, ELISAs were performed essentially as above, except that wells were coated with 10µg/ml anti-mouse IgG (Fab-specific, Sigma, Poole, UK) 5µg/ml purified F1-antigen in PBS or 6µg/ml *S. typhimurium* SL3261 lysate. Purified IgG₁ or IgG_{2a} (Sigma, Poole, UK) and day 98 serum
10 samples were diluted in BLOTTO. Peroxidase-labelled secondary antibodies against mouse IgG₁ or IgG_{2a} were diluted 1:4000 BLOTTO before use.

- The results (Fig 5) indicated that SL3261 containing pP_{phoP}-F1, pP_{lacZ}-F1 or pP_{pagC}-F1 plasmids all induced serum IgA antibody to
15 F-antigen. The induction of circulating IgA to F1-antigen did not correlate with the presence of IgA to F1-antigen at mucosal surfaces. For example, SL3261 / pP_{lacZ}-F1 induced high levels of serum antibody to F1-antigen but IgA antibody to F1-antigen was
20 not detected in gut or lung wash samples. Only SL3261 / pP_{phoP}-F1 induced an IgA antibody response to F1-antigen in both the gut and the lung.

Claims

1. A recombinant gut-colonising microorganism which comprises a promoter selected from the P_{ompC} , P_{phoP} and P_{pagC} or
5 fragments or variants thereof which can act as promoters, said promoter being operatively interconnected with a nucleic acid which encodes a heterologous protein, able to induce a protective immune response against a different organism, in a mammal to which it is administered.
 - 10 2. A recombinant gut-colonising microorganism according to claim 1 wherein the promoter is P_{phoP} promoter.
 3. A recombinant gut-colonising microorganism according to
15 claim 1 or claim 2 which comprises a *Salmonella* spp.
 4. A recombinant gut-colonising microorganism according to claim 3 wherein the *Salmonella* spp. is *Salmonella typhimurium* or *Salmonella typhi*.
 - 20 5. A recombinant gut-colonising microorganism according to any one of the preceding claims comprising wherein the gut-colonising microorganism is attenuated.
 - 25 6. A recombinant gut-colonising microorganism according to any one of the preceding claims wherein the heterologous protein, is able to induce a protective immune response against *Yersinia pestis*.
 - 30 7. A recombinant gut-colonising microorganism according to claim 6 wherein the said heterologous protein comprises an F1-antigen of *Yersinia pestis* or an antigenic fragment or variant thereof.
 - 35 8. A vaccine comprising a recombinant gut-colonising microorganism according to any one of the preceding claims.
-
-

9. A vaccine according to claim 8 which further comprises a pharmaceutically acceptable carrier or diluent.

10. A vaccine according to claim 8 or claim 9 which is
5 adapted for oral administration.

11. A method of inducing a protective immune response against a pathogen in a mammal, said method comprising administering to said mammal a recombinant gut-colonising
10 microorganism which comprises a promoter selected from the P_{ompC} , P_{phoP} and P_{pagC} or fragments or variants thereof which can act as promoters, said promoter being operatively interconnected with a nucleic acid which encodes an antigen protein, able to induce a protective immune response against said pathogen, in a mammal to
15 which it is administered.

12. The use of a promoter selected from P_{ompC} , P_{phoP} and P_{pagC} in the production of a vaccine comprising a recombinant gut-colonising organism.

Abstract

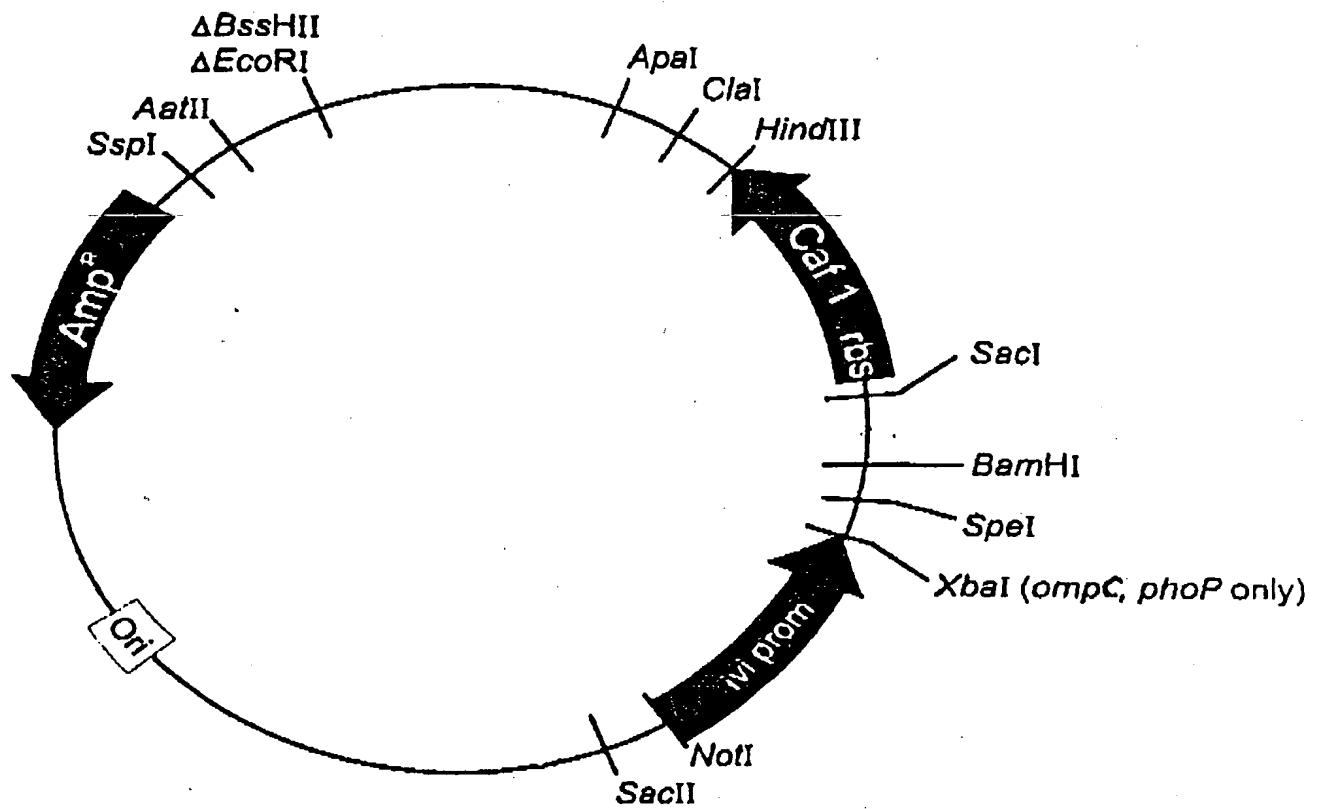
A recombinant gut-colonising microorganism such as a *Salmonella* spp. which comprises a promoter selected from the P_{ompC} , P_{phoP} and
5 P_{pagC} or fragments or variants thereof which can act as promoters, said promoter being operatively interconnected with a nucleic acid which encodes a heterologous protein, able to induce a protective immune response against a different organism, in a mammal to which it is administered.

10

Such organisms are useful in the preparation of vaccines.

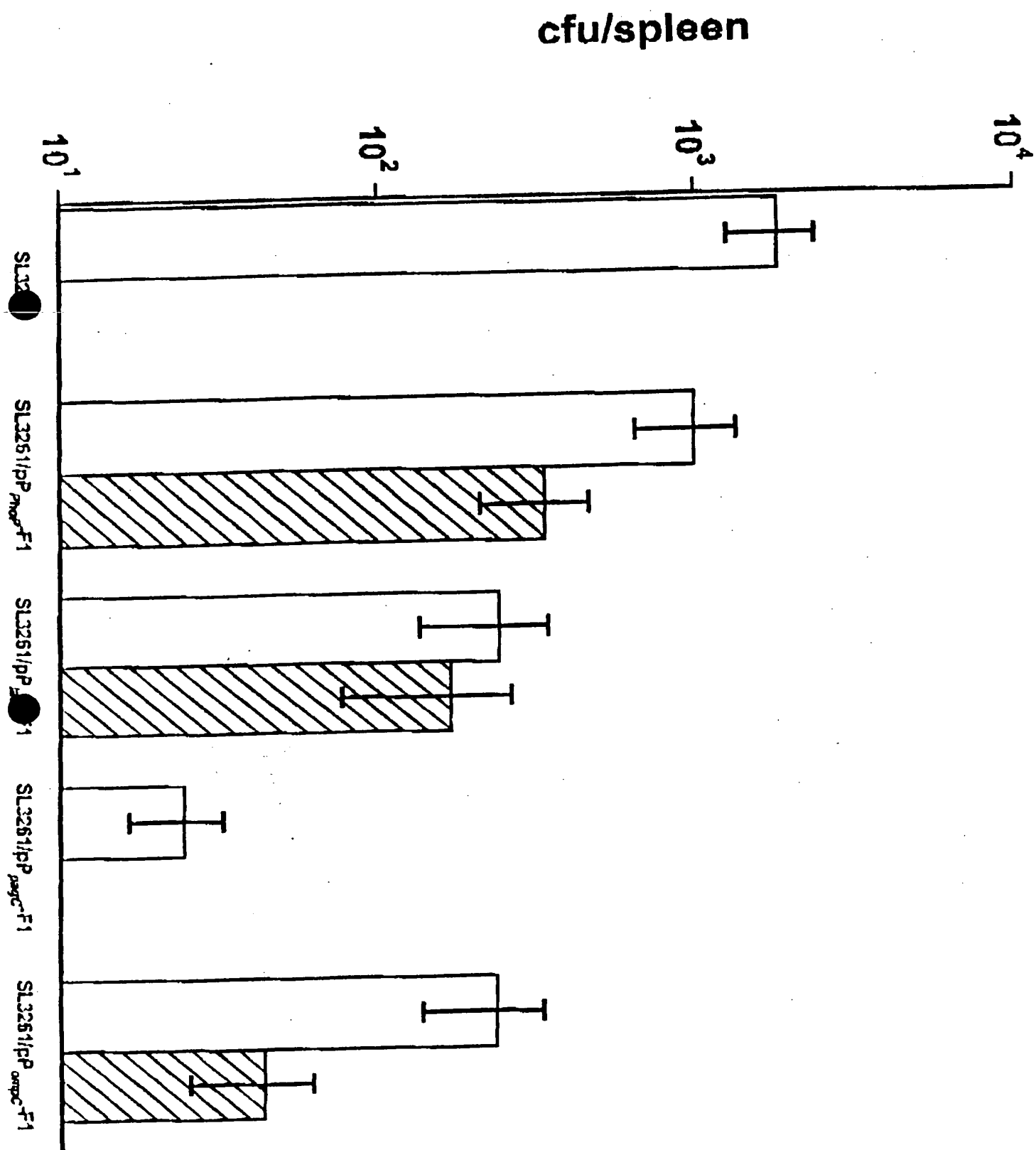
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Figure 1



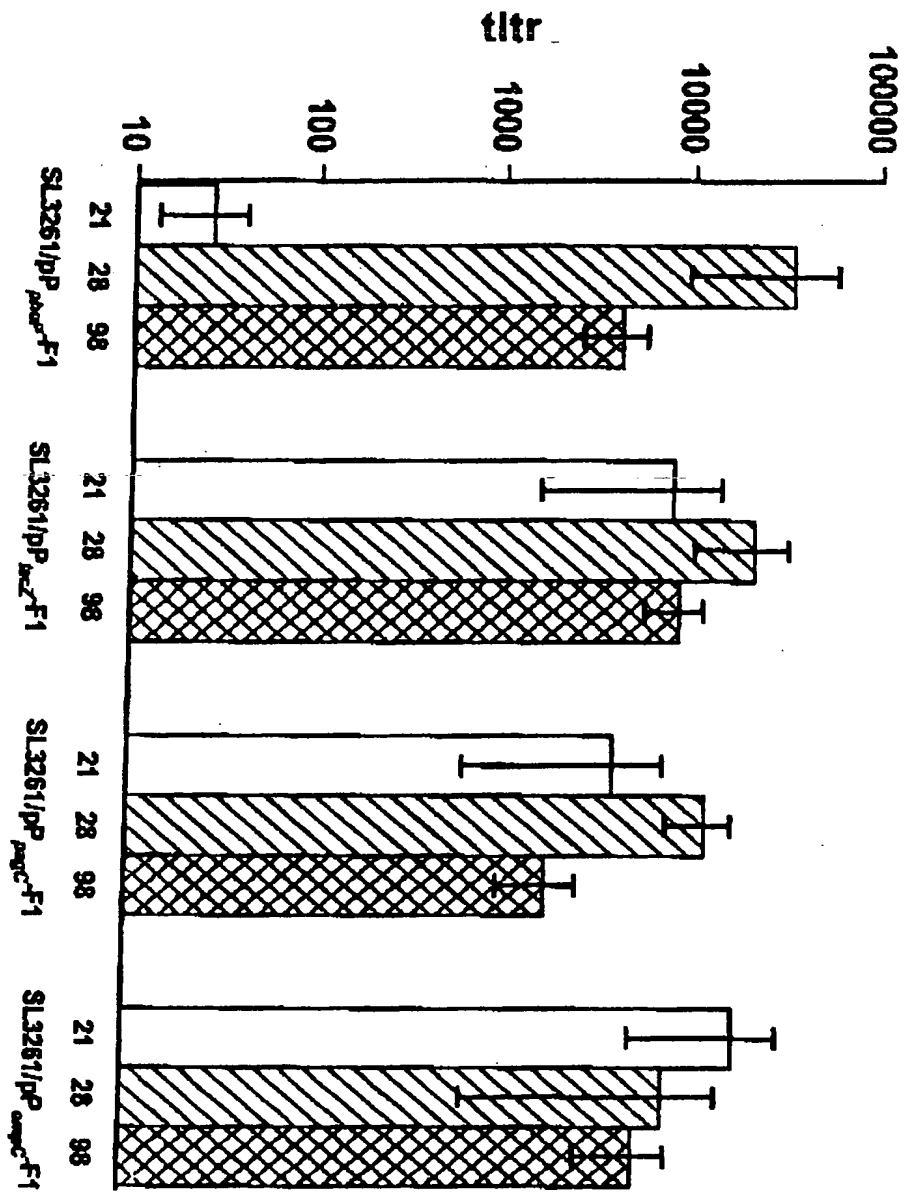
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Figure 2



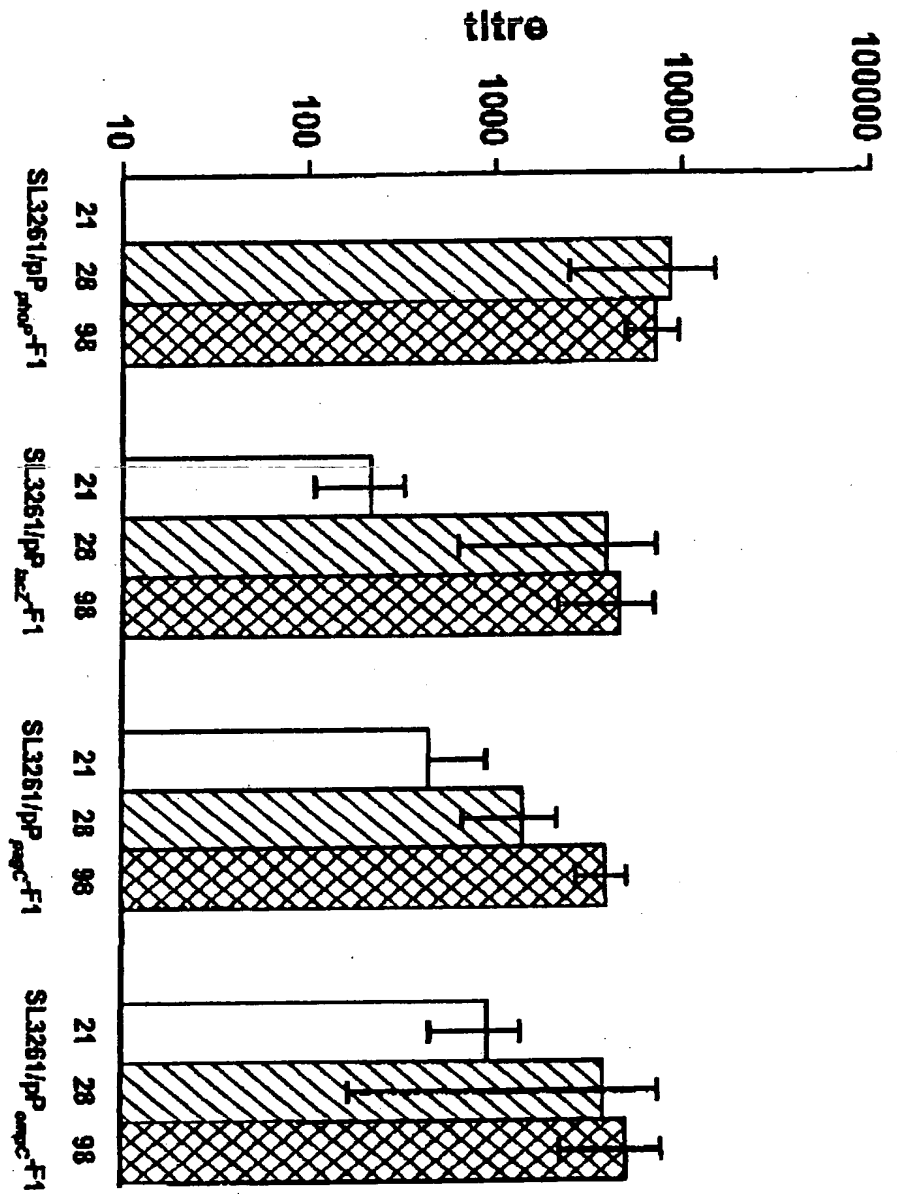
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Figure 3a



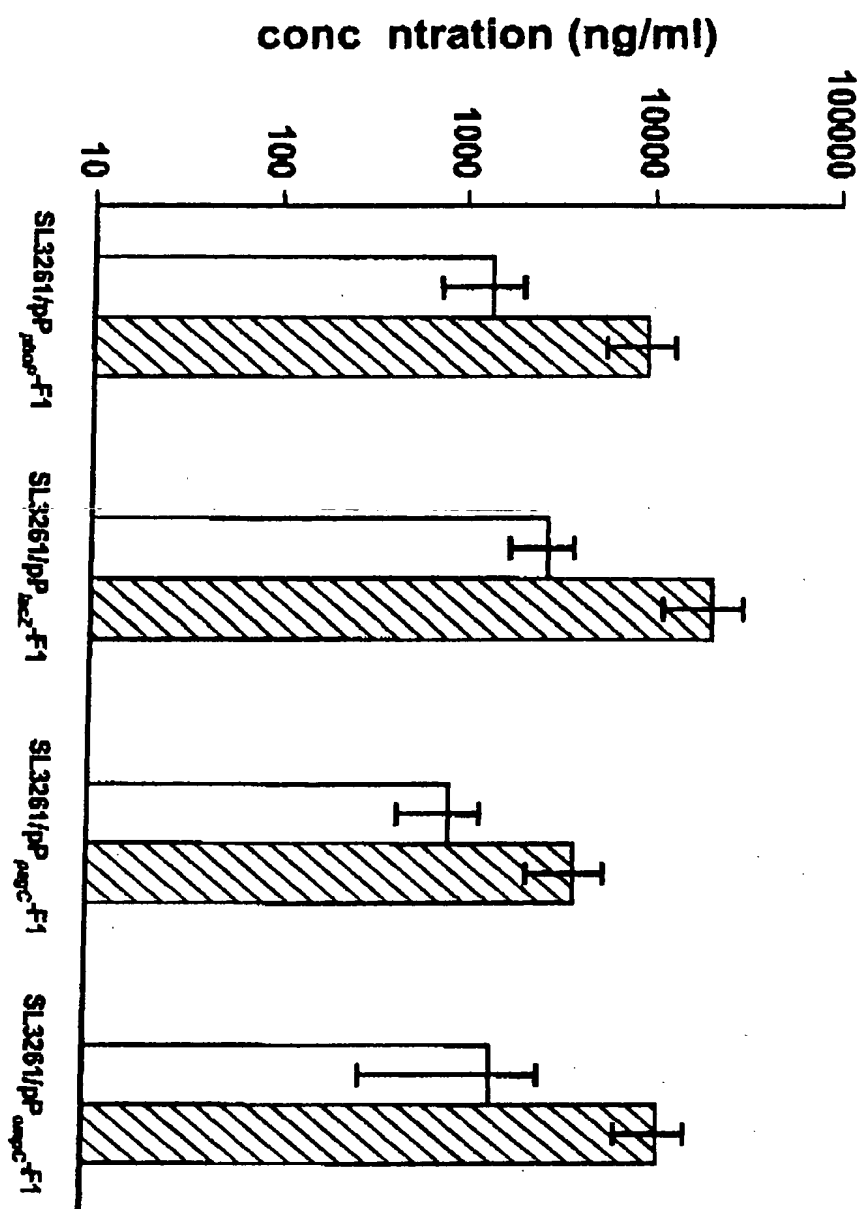
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Figure 3b



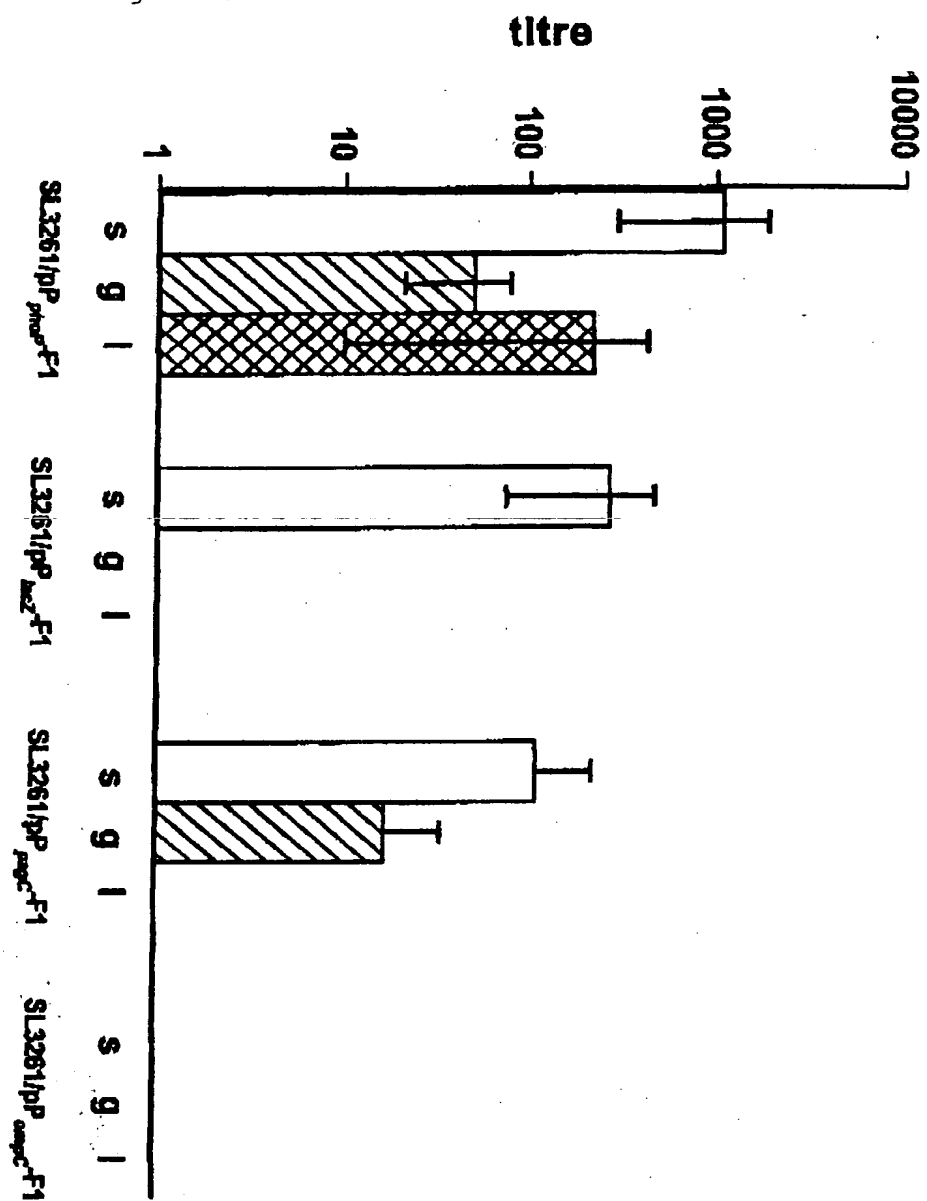
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Figure 4



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Figure 5



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Figure 6

P_{lac} 5'
CGCCCAATACGCAAACCGCCTCTCCCCGCGCGTTGGCCGATTCATTAATGCAGCTGGCACGACAGGTTTCC
CGACTGGAAAGCGGGCAGTGAGCGCAACGCAATTAATGTGAGTTAGCTCACTCATTAGGCACCCCAGGCTT
TACACTTTATGCTTCCGGCTCGTATGTTGTGTGGAATTGTGAGCGGATAACAAT 3'
(SEQ ID NO 1)

P_{phoP} 5'
GTGACTCTGGTCGACGAACTTAAATAATGCCTGCCTCACCTCTTTTCTTCAGAAAGAGGGTGACTATTTG
TCTGGTTTATTAAGTGTATCCCAAAGCACCATAATCAACGCTAGACTGTTCTTATTGTTAACACA 3'
(SEQ ID NO 2)

P_{pagC} 5'
GTTAACCCTCTTAATAATAATGGGTTTTATAGCGAAATAGACTTTTTATCGCGTGTTCAATATTTGCGT
TAGTTATTATTTTTTTGGAATGTAAATTCTCTCTAAACACAGGTGATATTTATGTTGGAATTGTGGTGTG
ATTCTATTCTTATAATATAACAAGAAATGTTGTAAGTATAGATATATTAAAAGATTAAATCGGAGCGGGA
ATAAAGCGTGCTAAGCATCATCGTGAATATGATTACAGCGCCTGCGATGGCATATAACCGTATTGCGGATG
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ATGGGTGATTTGTTGTTCTATAGTGGCTAAACACTTTATGGTTTCTGTTAAATATATATGCGTGAGAAAAA
TTAGCATTCAAATCTATAAAAGTTAGATGACATTGTAGAACCGGTTACCTAAATGAGCGATAGAGTGCTTC
GGTAGTAAAAATATCTTTCAGGAAGTAAACACATCAGGAGCGATAGCGGTGAATTATTCGTGGTTTTGTCC
ATTCGGCATAGTGGCGATAACTGAATGCCGGATCGGTACTGCAGGTGTTTAAACACACCGTAAATAATAAG
TAGTA 3'
(SEQ ID NO 3)

P_{ompC} 5'
TAAACAGACATTCAGAAGTGAATGACGGTAATAAAATAAAGTTAATGATGATAGCGGTCACTATTTTAGTTG
CGAATGAAGATTCTGTTTTATCATTCACTGCTATGAATTTTCATCAATTTAACCCGTTGATTTTAAAAGTTT
CGTGAAATATATTTTGCTATTTGTGCTTATTTTTACTTGATTTTGGCTTTAAAAAAGTTCCGTAAATTC
ATATTTTGAAACATCTATATAGATAACTGTAACATCTTAAAGTTTTAGTATCATATTCGTGTTGGATTAT
TCTGTATTTTGGCGAGAATGGACTTGCCGACTGGTTAATGAGGGTTAACCAGTAAGCAGTGGCATAAAAA
AGCAATAAAGGCATAT 3'
(SEQ ID NO 4)

The locations of oligonucleotide primers used in the PCR are shown underlined. Promoter regions are shown in bold and repeated sequences upstream of these promoters are shown double underlined

PCT 1 B00103402

(Tracey Carter)

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